**ICBF Beef Industry consultation meeting held on Tuesday 21st June 2016 from 7pm to 9pm at the Killeshin Hotel, Portlaoise, Co. Laois.**

**Attending:**

Sean Coughlan, Andrew Cromie, Stephen Conroy, John Appleby, Tom Mulligan, Gerry Kilgallon, Nevan McKiernan, Alan Wood, Eamonn Coleman, Mary Duggan, Meryl Gillespie, Richard Fortune, Joe Fortune, John Burke, Matt McClure, Jim Parkinson, Gerard Dineen, Francis Kearney, Ciaran Costello, Denis Large, Patrick O’ Donnell, John Clarke, Michael Conway, Mary Purcell, Michael McKeon, Peter Luttrell, Mareille McCall, Alfie Hanbidge, David Clarke, John Lynch, Garrett Behan, Peadar Glennon, Roger O’ Halloran, Larry Feeney, John McEnroe, Louis Byrne, Paul Sykes, Bernard Eivers, Doreen Corridan, Gerry Greally, Hohn Carty, Leslie Sandes, Margaret Kelleher, Michael Doran, Noirin McHugh, Robert Prendiville, Stephen Conroy, Eddie Brennan, David Johnson, David Holland, Lisa Ring, Ross Evans, Hugh Farrell, Jimmy Cosgrave, Karl O’ Connell, Tom Wilson, Michael Gunn, Shane Beirne, Sean Sherman, Stephen Connolly, Nicola Hobson, Martin Ryan, Kieran Mailey, Shane Murphy, Laurie Harney, Aidan Murray & Chris Daly.

**Apologies:** Joe Deverell, Ger Ryan, Anselm Fitzgerald.

# 1. Beef.

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| **Topic** | **Key points** | **Actions/Decisions** |
| Validation of €uro-Star Replacement Index – Donagh Berry. | * Analysis was shown on 162,363 females that were born in 2011 and subsequently kept as replacements in the suckler herd. Overall the analysis showed that the 5 star females outperformed the 1 star females and left more profit to the farmer. Also, when genomic data was included in the replacement index, it was a more accurate predictor of their subsequent performance when compared to replacement index of the group based on parent average only. | Further validation work is to be carried out and presented at future meetings. |
| Beef genomic evaluations – Ross Evans | * A number of changes were made to the beef genomic AI bull test proof evaluations since the last meeting. The latest AI test proofs with the changes included were circulated prior to the meeting. These changes were as follows:  1. New phenotypic data and pedigree included up to the 16th May 2016. 2. Survival trait is now evaluated using a two-step process similar to dairy genomic evaluations. 3. 418 thousand beef genotypes in the evaluation | * Include the young AI bulls that were not in the test proof file in future versions that will be circulated. * Further investigate traits such as mortality and cow live-weight which are increasing with genomics. * Carry out further validation work where possible on the beef genomic evaluations. This validation work will include sending out €uro-Star reports on breeding females to a number of pedigree and commercial performance recording herds with both the current replacement index (without genomics included) and the replacement index with genomics included for feedback. * Final set of genomic test proofs will be available in advance of the next meeting, scheduled for the2nd August (evening meeting). |